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RAW SEQUENCE LISTING

DATE: 06/25/2003

PATENT APPLICATION: US/09/804,409B

TIME: 16:19:06

Input Set : A:\0278721.APP.txt

Output Set: N:\CRF4\06252003\I804409B.raw

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SEP 22 2003
TECH CENTER 1600/2900

3 <110> APPLICANT: ENGINE, INC.
4 KIEFFER, TIMOTHY J.
5 CHEUNG, ANTHONY T.
7 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR REGULATED PROTEIN
8 EXPRESSION IN GUT
10 <130> FILE REFERENCE: 029996/027 8721
12 <140> CURRENT APPLICATION NUMBER: 09/804,409B
13 <141> CURRENT FILING DATE: 2001-03-12
15 <160> NUMBER OF SEQ ID NOS: 19
17 <170> SOFTWARE: PatentIn Ver. 2.1
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24 <220> FEATURE:
25 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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60 <220> FEATURE:
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68 <211> LENGTH: 1319

69 <212> TYPE: DNA

70 <213> ORGANISM: Mus musculus

71 <400> SEQUENCE: 5

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75 gtttgttgaa tgaatacacg cgaagccggt tctcatttag gggcatgagt aggcagaggt 180
76 gtgggcagga agcaggaaaag agcggaaaca ggtgoggaca gaaaggaggg gctctgaagg 240
77 atgcacgtca gtgcacaaat gtcctccaga taccaggttc actgtggccc taggccaggc 300
78 tgcacggggc tcccatgtg gtctgcccag ggtgagagca gaactgggtt gggcggggca 360
79 gaagaaaacc aaccaggaaag cagggttgca cccaaattat ccaggtttta agtacattta 420
80 agagacaagg ctgggctgtt gaaggtcaga ggtgtccctg ggggtgtgga ctaggactga 480
81 ccactctgtt tttagtttaa tgggtgagaa tgcctcacac tgcctcctgc cttacttgcc 540
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84 cactatccc talcaataaa caattaaata cacacagaat gogaggcaca caactgagtt 720
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92 agccggcata gaattgctgc tgcctgcggc gccacggcca ccatcacggc tgttaccacc 1200
93 accgtaactg cagtgttccc gctgggtcag agctttggta gccagactac agaccactc 1260
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100 <213> ORGANISM: Mus musculus

101 <400> SEQUENCE: 6

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105 gtcttgctag caaaaatttt ctggcatatg caatagtgtc tgggttttgt ggttgtatat 180
106 gggctggatc cccgggtggg gcagtctctg gatggctctt ccttcogtct tagctccaaa 240
107 ctttgtctct gtaactcctt ccattgggtac tttgtttccc attctaagaa ggagcaaaat 300
108 atccacaatt cctttctctt cctttctctt gagttttgca aatgcacaaa aactttcaaa 360
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112 aattggaatg tgattgttca gtcagcagag acttttagag gaacctatcc aacaagattc 600
113 tctcagttct cagaaaatata ttccagtata tacagggtta gaggactcac atctttaata 660
114 aaataaaagt aaaaatttag acctgtataa attattaagg tacctaatac agttccacgg 720
115 caaagtacag ccattggtat gaattataaa tccaagaagc ggtgggttaa ctctgacatt 780
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119 aatccctccc ttctggtag gcagtatgtt ttttgagca cagtttctta gctatctctt 1020
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122 aaaaaattaa gcatctcacc ttttttgctc aaactaaaac gttttaaaac agttctgcoct 1100
123 ggagtcatga tatgaaatac gatctatcat atttgcaatg ttctgttcaa ttgtggctgc 1260
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125 aaattcttat gacatggcag caagcccaag aaacctttct aaacaaggcg tgaaaacgca 1380
126 gagatgtcct tgcgaattag catgtctatc tgacagattt ctccctttct aagggaattt 1440
127 gtgctgaaca ttttatttct agcctcagag ataaaagaag ggggaagaag ctgtagtttt 1500
128 tgctacataa gacagggtgg gtaagcatgc aacgctttta aaaaatatct aaagtgattg 1560
129 tttctctctg gattctttga aaaagctcgc ctgogctggg gtttgagggt gagccggtga 1620
130 cgtcagcgtg gaatggggag tcaggcgccc aggcctctct taagccgagg agctgtcccg 1680
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136 <13> ORGANISM: Mus musculus
137 <40> SEQUENCE: 7
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143 tggcctacca cagatttcat gtctggcaact ggctatgtca gaacatgtag gagcttttgg 180
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171 aaaaaaata aaaaataaaa tattagaata aaatgtagag gaatattttt aatttaacaa 1860
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176 ttcaatagta atataattat tgaacaaata atccttaaaa gaagaaatcc agagggaatag 1510
177 caagtttaggg gaagagagggg tgtgtgtgtg tgtgtgtgtg cgcacattta tagccaaaat 1520
178 agatgatata ctttaattgaa catgccaatta aaaccacatta ttttgcatac agtttacata 1530
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181 tctttctgta tctctgaaga aagtgaacgt gtacttataa ttataaataa ataaatcttt 1560
182 aacccaaaaa ccccccataat ttcaacaaac gatatgtcct ggtctgaggg ttccagggcat 1570
183 agaaatagaa acacacagag tgtggagaca gtgoggttca ggtccggcat tccagttcag 1580
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187 <110> SEQ ID NO: 8

188 <111> LENGTH: 226

189 <112> TYPE: DNA

190 <113> ORGANISM: Homo sapiens

191 <400> SEQUENCE: 8

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194 cggcgggggg ccagcgtggc ccaggggaaa gcagaggggg caccgagcgg gcagagaccc 180
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200 <111> LENGTH: 110

201 <112> TYPE: PRT

202 <113> ORGANISM: Homo sapiens

203 <400> SEQUENCE: 9

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207      20          25          30
208 Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe
209      35          40          45
210 Phe Tyr Thr Pro Lys Thr Arg Arg Glu Ala Glu Asp Leu Gln Val Gly
211      50          55          60
212 Gln Val Glu Leu Gly Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu
213      65          70          75          80
214 Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys
215      85          90          95
216 Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn
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225 <112> TYPE: DNA

226 <113> ORGANISM: Homo sapiens

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230 tgaaccaaca cctgtgcggc tcacaccgg tggaagctct ctacctagtg tgggggggaa 180
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238 tgcagaagcg tggcattgtg gaacaatgct gtaccagcat ctgctccctc taccagctgg 360
239 agaactactg caactagaag cagcccgag gcagccccc acccgccgoc toctgcaocg 420
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244 <211> LENGTH: 167

245 <212> TYPE: PRT

246 <213> ORGANISM: Homo sapiens

248 <400> SEQUENCE: 11

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252             20             25             30
253 Thr Leu Ile Lys Thr Ile Val Thr Arg Ile Asn Asp Ile Ser His Thr
254             35             40             45
255 Gln Ser Val Ser Ser Lys Gln Lys Val Thr Gly Leu Asp Phe Ile Pro
256             50             55             60
257 Gly Leu His Pro Ile Leu Thr Leu Ser Lys Met Asp Gln Thr Leu Ala
258             65             70             75             80
259 Val Tyr Gln Gln Ile Leu Thr Ser Met Pro Ser Arg Asn Val Ile Gln
260             85             90             95
261 Ile Ser Asn Asp Leu Glu Asn Leu Arg Asp Leu Leu His Val Leu Ala
262             100            105            110
263 Phe Ser Lys Ser Cys His Leu Pro Trp Ala Ser Gly Leu Glu Thr Leu
264             115            120            125
265 Asp Ser Leu Gly Gly Val Leu Glu Ala Ser Gly Tyr Ser Thr Glu Val
266             130            135            140
267 Val Ala Leu Ser Arg Leu Gln Gly Ser Leu Gln Asp Met Leu Trp Gln
268             145            150            155            160
269 Leu Asp Leu Ser Pro Gly Cys

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280 165

283 <210> SEQ ID NO: 12

284 <211> LENGTH: 3408

285 <212> TYPE: DNA

286 <213> ORGANISM: Homo sapiens

288 <400> SEQUENCE: 12

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291 caagatgaca ccaaaacct catcaagaca attgtcacca ggatcaatga catttcacac 180
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293 cccatcttga ccttatccaa gatggaccag acaactggcag tctacccaca gatctccacc 300
294 agtatgcctt ccagaaaagt gatccaaata tccaaacgacc tggagaaact ccgggatctt 360
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